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SEP 10 2003
TECH CENTER 1600

RAW SEQUENCE LISTING DATE: 09/05/2003
PATENT APPLICATION: **US/10/081,817A** TIME: 06:56:07

Input Set : **A:\00530-094001.txt**
Output Set: **N:\CRF4\09052003\J081817A.raw**

<110> APPLICANT: Polyak, Kornelia
Porter, Dale
Sgroi, Dennis
Krop, Ian
<120> TITLE OF INVENTION: HIN-1, A TUMOR SUPPRESSOR GENE
<130> FILE REFERENCE: 00530-094001
<140> CURRENT APPLICATION NUMBER: US 10/081,817A
<141> CURRENT FILING DATE: 2002-02-22
<150> PRIOR APPLICATION NUMBER: US 60/270,973
<151> PRIOR FILING DATE: 2001-02-23
<150> PRIOR APPLICATION NUMBER: US 60/351,908
<151> PRIOR FILING DATE: 2002-01-25
<160> NUMBER OF SEQ ID NOS: 32
<170> SOFTWARE: FastSEQ for Windows Version 4.0

ENTERED

<210> SEQ ID NO: 1
<211> LENGTH: 104
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens
<400> SEQUENCE: 1

```
Met Lys Leu Ala Ala Leu Leu Gly Leu Cys Val Ala Leu Ser Cys Ser
 1               5                  10                  15
Ser Ala Arg Ala Phe Leu Val Gly Ser Ala Lys Pro Val Ala Gln Pro
            20                  25                  30
Val Ala Ala Leu Glu Ser Ala Ala Glu Ala Gly Ala Gly Thr Leu Ala
        35                  40                  45
Asn Pro Leu Gly Thr Leu Asn Pro Leu Lys Leu Leu Leu Ser Ser Leu
    50                  55                  60
Gly Ile Pro Val Asn His Leu Ile Glu Gly Ser Gln Lys Cys Val Ala
65                  70                  75                  80
Glu Leu Gly Pro Gln Ala Val Gly Ala Val Lys Ala Leu Lys Ala Leu
                85                  90                  95
Leu Gly Ala Leu Thr Val Phe Gly
            100
```

<210> SEQ ID NO: 2
<211> LENGTH: 86
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens
<400> SEQUENCE: 2

```
Arg Ala Phe Leu Val Gly Ser Ala Lys Pro Val Ala Gln Pro Val Ala
 1               5                  10                  15
Ala Leu Glu Ser Ala Ala Glu Ala Gly Ala Gly Thr Leu Ala Asn Pro
            20                  25                  30
Leu Gly Thr Leu Asn Pro Leu Lys Leu Leu Leu Ser Ser Leu Gly Ile
```

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```
58         35                  40                  45
59 Pro Val Asn His Leu Ile Glu Gly Ser Gln Lys Cys Val Ala Glu Leu
60     50                  55                  60
61 Gly Pro Gln Ala Val Gly Ala Val Lys Ala Leu Lys Ala Leu Leu Gly
62 65                  70                  75                  80
63 Ala Leu Thr Val Phe Gly
64                 85
66 <210> SEQ ID NO: 3
67 <211> LENGTH: 312
68 <212> TYPE: DNA
69 <213> ORGANISM: Homo sapiens
71 <400> SEQUENCE: 3
72 atgaagctcg ccgccctcct ggggctctgc gtggccctgt cctgcagctc cgctcgtgct      60
73 ttcttagtgg gctcggccaa gcctgtggcc cagcctgtcg ctgcgctgga gtcggcggcg     120
74 gaggccgggg ccgggaccct ggccaacccc ctcggcaccc tcaacccgct gaagctcctg     180
75 ctgagcagcc tgggcatccc cgtgaaccac ctcatagagg gctcccagaa gtgtgtggct     240
76 gagctgggtc cccaggccgt gggggccgtg aaggccctga aggccctgct gggggccctg     300
77 acagtgtttg gc                                                        312
79 <210> SEQ ID NO: 4
80 <211> LENGTH: 258
81 <212> TYPE: DNA
82 <213> ORGANISM: Homo sapiens
84 <400> SEQUENCE: 4
85 cgtgctttct tagtgggctc ggccaagcct gtggcccagc ctgtcgctgc gctggagtcg      60
86 gcggcggagg ccggggccgg gaccctggcc aacccccctcg gcaccctcaa cccgctgaag     120
87 ctcctgctga gcagcctggg catccccgtg aaccacctca tagagggctc ccagaagtgt     180
88 gtggctgagc tgggtcccca ggccgtgggg gccgtgaagg ccctgaaggc cctgctgggg     240
89 gccctgacag tgtttggc                                                  258
91 <210> SEQ ID NO: 5
92 <211> LENGTH: 104
93 <212> TYPE: PRT
94 <213> ORGANISM: Mus musculus
96 <400> SEQUENCE: 5
97 Met Lys Leu Thr Thr Thr Phe Leu Val Leu Cys Val Ala Leu Leu Ser
98  1               5                  10                  15
99 Asp Ser Gly Val Ala Phe Phe Met Asp Ser Leu Ala Lys Pro Ala Val
100             20                  25                  30
101 Glu Pro Val Ala Ala Leu Ala Pro Ala Ala Glu Ala Val Ala Gly Ala
102         35                  40                  45
103 Val Pro Ser Leu Pro Leu Ser His Leu Ala Ile Leu Arg Phe Ile Leu
104     50                  55                  60
105 Ala Ser Met Gly Ile Pro Leu Asp Pro Leu Ile Glu Gly Ser Arg Lys
106 65                  70                  75                  80
107 Cys Val Thr Glu Leu Gly Pro Glu Ala Val Gly Ala Val Lys Ser Leu
108                 85                  90                  95
109 Leu Gly Val Leu Thr Met Phe Gly
110             100
112 <210> SEQ ID NO: 6
113 <211> LENGTH: 85
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<212> TYPE: PRT
<213> ORGANISM: Mus musculus
<400> SEQUENCE: 6

Val	Ala	Phe	Phe	Met	Asp	Ser	Leu	Ala	Lys	Pro	Ala	Val	Glu	Pro	Val
1				5				10					15		
Ala	Ala	Leu	Ala	Pro	Ala	Ala	Glu	Ala	Val	Ala	Gly	Ala	Val	Pro	Ser
		20					25					30			
Leu	Pro	Leu	Ser	His	Leu	Ala	Ile	Leu	Arg	Phe	Ile	Leu	Ala	Ser	Met
	35					40					45				
Gly	Ile	Pro	Leu	Asp	Pro	Leu	Ile	Glu	Gly	Ser	Arg	Lys	Cys	Val	Thr
50					55					60					
Glu	Leu	Gly	Pro	Glu	Ala	Val	Gly	Ala	Val	Lys	Ser	Leu	Leu	Gly	Val
65				70					75					80	
Leu	Thr	Met	Phe	Gly											
				85											

<210> SEQ ID NO: 7
<211> LENGTH: 312
<212> TYPE: DNA
<213> ORGANISM: Mus musculus
<400> SEQUENCE: 7

atgaagctta ccaccacctt tctagtgctc tgtgtggctc tgctcagtga ctctggtgtt 60
gctttcttca tggactcatt ggccaagcct gcggtagaac ccgtggccgc ccttgctcca 120
gctgcagagg ctgtggcagg ggctgtgcct agcctaccat taagccactt ggccatcctg 180
aggttcatcc tggccagcat gggcatccca ttggatcctc tcatagaggg atccaggaag 240
tgtgtcaccg agctgggccc tgaggctgta ggagctgtga agtcactgct gggggtcctg 300
acaatgttcg gt 312

<210> SEQ ID NO: 8
<211> LENGTH: 255
<212> TYPE: DNA
<213> ORGANISM: Mus musculus
<400> SEQUENCE: 8

gttgctttct tcatggactc attggccaag cctgcggtag aacccgtggc cgcccttgct 60
ccagctgcag aggctgtggc aggggctgtg cctagcctac cattaagcca cttggccatc 120
ctgaggttca tcctggccag catgggcatc ccattggatc ctctcataga gggatccagg 180
aagtgtgtca ccgagctggg ccctgaggct gtaggagctg tgaagtcact gctgggggtc 240
ctgacaatgt tcggt 255

<210> SEQ ID NO: 9
<211> LENGTH: 23
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: primer
<400> SEQUENCE: 9

gagggaaagt ttttttttatt tgg 23

<210> SEQ ID NO: 10
<211> LENGTH: 22
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:

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Input Set : **A:\00530-094001.txt**
Output Set: **N:\CRF4\09052003\J081817A.raw**

173 <223> OTHER INFORMATION: primer
175 <400> SEQUENCE: 10
176 caaaactaac aaaacaaaac ca 22
178 <210> SEQ ID NO: 11
179 <211> LENGTH: 24
180 <212> TYPE: DNA
181 <213> ORGANISM: Artificial Sequence
183 <220> FEATURE:
184 <223> OTHER INFORMATION: primer
186 <400> SEQUENCE: 11
187 gttaagagga agttttcgag gttc 24
189 <210> SEQ ID NO: 12
190 <211> LENGTH: 24
191 <212> TYPE: DNA
192 <213> ORGANISM: Artificial Sequence
194 <220> FEATURE:
195 <223> OTHER INFORMATION: primer
197 <400> SEQUENCE: 12
198 ggtacgggtt ttttacggtt cgtc 24
200 <210> SEQ ID NO: 13
201 <211> LENGTH: 22
202 <212> TYPE: DNA
203 <213> ORGANISM: Artificial Sequence
205 <220> FEATURE:
206 <223> OTHER INFORMATION: primer
208 <400> SEQUENCE: 13
209 aacttcttat acccgatcct cg 22
211 <210> SEQ ID NO: 14
212 <211> LENGTH: 24
213 <212> TYPE: DNA
214 <213> ORGANISM: Artificial Sequence
216 <220> FEATURE:
217 <223> OTHER INFORMATION: primer
219 <400> SEQUENCE: 14
220 gttaagagga agtttttgag gttt 24
222 <210> SEQ ID NO: 15
223 <211> LENGTH: 24
224 <212> TYPE: DNA
225 <213> ORGANISM: Artificial Sequence
227 <220> FEATURE:
228 <223> OTHER INFORMATION: primer
230 <400> SEQUENCE: 15
231 ggtatgggtt ttttatggtt tgtt 24
233 <210> SEQ ID NO: 16
234 <211> LENGTH: 25
235 <212> TYPE: DNA
236 <213> ORGANISM: Artificial Sequence
238 <220> FEATURE:
239 <223> OTHER INFORMATION: primer

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241 <400> SEQUENCE: 16
242 caaaacttct tatacccaat cctca 25
244 <210> SEQ ID NO: 17
245 <211> LENGTH: 21
246 <212> TYPE: DNA
247 <213> ORGANISM: Artificial Sequence
249 <220> FEATURE:
250 <223> OTHER INFORMATION: primer
252 <400> SEQUENCE: 17
253 tttccctgct tccacactag c 21
255 <210> SEQ ID NO: 18
256 <211> LENGTH: 21
257 <212> TYPE: DNA
258 <213> ORGANISM: Artificial Sequence
260 <220> FEATURE:
261 <223> OTHER INFORMATION: primer
263 <400> SEQUENCE: 18
264 agattaagaa ggaattgacc t 21
266 <210> SEQ ID NO: 19
267 <211> LENGTH: 551
268 <212> TYPE: DNA
269 <213> ORGANISM: Homo sapiens
271 <220> FEATURE:
272 <221> NAME/KEY: misc_feature
273 <222> LOCATION: 189
274 <223> OTHER INFORMATION: n = C or G
276 <400> SEQUENCE: 19
277 cggccgggga ggcggccggg agtgaggcct gatcgtccct ggcgcctcca cctccccagg 60
278 cgcagaaggc gcccacgagg acccccagtg cccgacgttg ccacggtctg ggatcagagg 120
279 cagggaccag ggagccagga actgcgccgc ccccgccccc gccctggcgc gagggaagct 180
W--> 280 **ccctcaccng agggaagctc ccctcacccg gcccagccct gcaggggggc gcgtggggtc** 240
281 agaccgcaaa gcgaaggtgc gggccggggt gggcctcgcg gagacaaagg ccgggcctgc 300
282 ctctctcaga gggccccagc gcctgccaag aggaagtcct cgaggcccgg gcagggaagg 360
283 gggcacgggc ttcccagggc ccgccggccg cagcaggaag ttggccaggg cacggccgtg 420
284 agcggagcgg gcagggcttt ctcaggagcg cgggcgaggc cggcgctgga ggggcgagga 480
285 ccgggtataa gaagcctcgt ggccttgccc gggcagccgc aggttccccg cgcgccccga 540
286 gcccccgcgc c 551
288 <210> SEQ ID NO: 20
289 <211> LENGTH: 279
290 <212> TYPE: DNA
291 <213> ORGANISM: Rattus norvegicus
293 <400> SEQUENCE: 20
294 gttctctgtt ttgtgttggt aggcgttgct ttcttggtgg attcactggc caagcctgtg 60
295 gtagaacccg tggctgccat tgctacagct gcagaggctg tggcaggggc tgtgcctagc 120
296 ctaccattaa gccacttggc catcctgagg ttcatcgtga ccagcctggg catcccattg 180
297 gatcctctca tagatggttc caggaagtgc gtcaccgagc tgggccctga ggctgtagga 240
298 gctgtgaagt cactgctggg ggccctgaca acgttcggt 279
300 <210> SEQ ID NO: 21
301 <211> LENGTH: 93

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 09/05/2003
PATENT APPLICATION: **US/10/081,817A** TIME: 06:56:08

Input Set : **A:\00530-094001.txt**
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Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:19; N Pos. 189